	n	Δ	Document ID	Title
Τ.	×	US A1	20040121346	Designing and construction of transcription template for synthesizing cell-free protein, and dilution batchtype method of synthesizing cell-free wheat germ protein by using the same
7	×	KR	KR 2000018290	Method for determining the volume and the location of Athe semi-active mount of a multi-degree-of-freedom system - NoAbstract
e e	×	JP	2000066665	Automatic composition apparatus for electronic musical instrument, performs Amelody and music based on melody data and melody characteristics data included in selected music template

L10 3/02/

	I	Ocument ID	Title
1	US	20050032086	Methods of RNA and protein
	A1		synthesis
	TTC	20050032078	Methods for the detection,
2	A1	20030032078	analysis and isolation of
	AI		nascent proteins
	IIC	20050009013	METHODS FOR THE DETECTION,
3	A1	20030003013	ANALYSIS AND ISOLATION OF
			NASCENT PROTEINS
4	US	20040235029	In vitro translation system
	A1		
	US	20040014071	Methods for the detection,
5	A1		analysis and isolation of
			nascent proteins
6	US	20030190643	Sequencing by mass
	A1_		spectrometry
	TIC	20030092031	Methods for the detection,
7	A1	_ , , , , , , , , , , , , , , , , , , ,	analysis and isolation of
			nascent proteins
8	US	20020132248	N-terminal and C-terminal
	A1		markers in nascent proteins
			Methods for the detection,
9	US	6306628 B1	analysis and isolation of
			Nascent proteins
10	IIC	6303337 B1	N-terminal and C-terminal
10	0.5	0000001 DI	markers in nascent proteins

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38306 S ENDO?/AU OR SAWASAKI?/AU OR OGASAWARA?/AU
L1
         30974 S MOTODA?/AU OR YABUKI?/AU OR KIGAWA?/AU OR YOKOYAMA?/AU
L2
         24311 S "TRANSCRIPTION TEMPLATE" OR "DNA TEMPLATE" OR (TEMPLATE (S) D
L3
         373335 S "POLMERASE CHAIN REACTION" OR PCR
L4
             21 S (1ST OR 2D OR 2ND OR 3D) (2A) PRIMER
L5
             59 S (1ST OR 2D OR 2ND OR 3D) (2A) PCR
L6
L7
            658 S (TWO-STEP OR 2-STEP) (2A) PCR
L8
          10143 S NESTED (2A) PCR
         61263 S AMPLIFICAT? (S) PCR
L9
L10
             1 S PROTEION (S) SYNTHES?
L11
         314249 S PROTEIN (S) SYNTHES?
           6363 S L11 (S) CELL-FREE
L12
             57 S L1 AND L12
L13
             51 S L2 AND L12
L14
              2 S L2 AND L7
L15
L16
              3 S L1 AND L7
         610752 S "POLYMERASE CHAIN REACTION" OR PCR
L17
             11 S L8 AND L1
L18
L19
              0 S L18 AND L2
L20
              3 S L12 AND L7
             2 DUP REM L20 (1 DUPLICATE REMOVED)
L21
             6 DUP REM L18 (5 DUPLICATES REMOVED)
L22
L23
             29 DUP REM L13 (28 DUPLICATES REMOVED)
             25 DUP REM L14 (26 DUPLICATES REMOVED)
L24
             2 DUP REM L15 (0 DUPLICATES REMOVED)
· L25
L26
              1 DUP REM L16 (2 DUPLICATES REMOVED)
L27
             0 S L23 AND L7
L28
             0 S L24 AND L7
L29
             0 S L23 AND L8
L30
             0 S L24 AND L8
L31
             3 S L17 AND L7 AND L12
              2 DUP REM L31 (1 DUPLICATE REMOVED)
L32
          1216 S "AFFINITY TAG" OR "PROTEIN TAG" OR 6-HIS OR "6-HIS TAG" OR "6
L33
L34
             0 S L23 AND L33
             0 S L24 AND L33
L35
             0 S L33 AND L17 AND L12
L36
L37
             7 S L33 AND L12
             3 DUP REM L37 (4 DUPLICATES REMOVED)
L38
             0 S L33 AND L8
L39
             0 S L33 AND L7
L40
             4 S L33 AND L3
L41
              3 DUP REM L41 (1 DUPLICATE REMOVED)
L42
           927 S (FIRST OR SECOND OR THIRD) (3A) PRIMER
L43
           907 S (FIRST OR SECOND OR THIRD) (3A) OLIGONUCLEOT?
L44
             0 S L33 AND L43
L45
             0 S L33 AND L44
L46
             4 S L43 AND L7
L47
             0 S L44 AND L7
L48
             3 DUP REM L47 (1 DUPLICATE REMOVED)
L49
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